S.Fig 1. Isolated mitochondrial fraction purity. Immunoblot for voltage dependent anion channel (mitochondria), protein disulfide isomerase (PDI; Endoplasmic reticulum), pan cadherin (plasma membrane) and alpha-tubulin (cytoplasm) to demonstrate purity of cellular fractions. n=1

S.Fig 2. Isolated mitochondrial proteome identifies electron transport chain protein expression changes with several that are improved or partially with withdrawal. (A) Oxidative phosphorylation pathway with an overlay of log fold expression quantification of differentially expressed proteins (DEP) from the isolated mitochondrial proteome dataset from C2C12 myotubes treated with 10mM ammonium acetate for 24 h (Am) with 24 h of ammonia removal (WD) compared to the untreated (UnT) myotubes. (B) Protein heatmap of the same oxidative phosphorylation pathway for the following comparisons: Am vs UnT, WD vs Am, and WD vs UnT. Red= increased expression, green= decreased expression. DEP significance was taken at p<0.05. All experiments were performed in n=3 biological replicates. Significance was calculated using an unpaired Student's t-test.

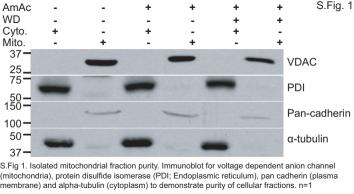
S.Fig 3. Unsupervised clustering reveals unique groups of molecules and their relationships across treatments. Heatmap with hierarchical clustering of rows and columns for A. Whole cell proteomics, B. Mitochondrial proteomics, and C. RNAseq from myotubes that are either untreated (UnT), treated for 24h with 10mM ammonium acetate (AmAc), and treated for 24h with AmAc and then have withdrawal (WD) of AmAc by media replacement for the final 24h of treatment. Dimensional reduction was performed using a p<0.05 for proteomics, and adjusted p-value of 0.05 for RNAseq using unpaired Student's t-test following Benjamini Hochberg correction. All experiments were performed in n=3 biological replicates.

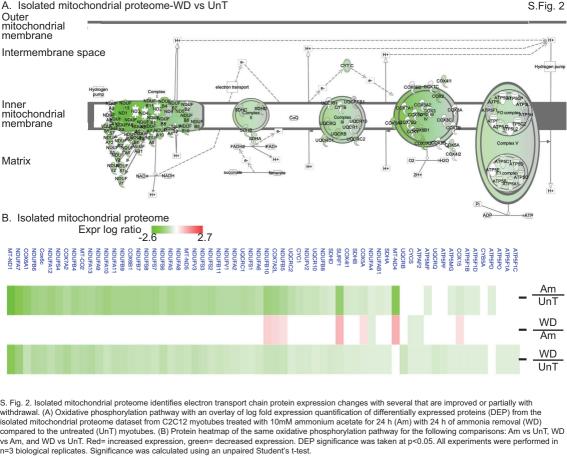
S.Fig 4. Hyperammonemia impairs respiration in mitochondria isolated from differentiated myotubes. Representative oxygraph tracings from isolated mitochondria from untreated (UnT) and 24 h of 10mM ammonium acetate (AmAc) treated differentiated myotubes. Oxygen consumption was measured in isolated mitochondria in respiration buffer in the basal state and in response to electron transport chain (ETC) complex substrates and inhibitors sequentially. After initial stabilization, 2 mM malate(M) and 2.5 mM pyruvate(P) were added. This was followed by 2.5 mM ADP(D); 10 mM glutamate(G); 10 mM succinate(S); 2 uM increments of

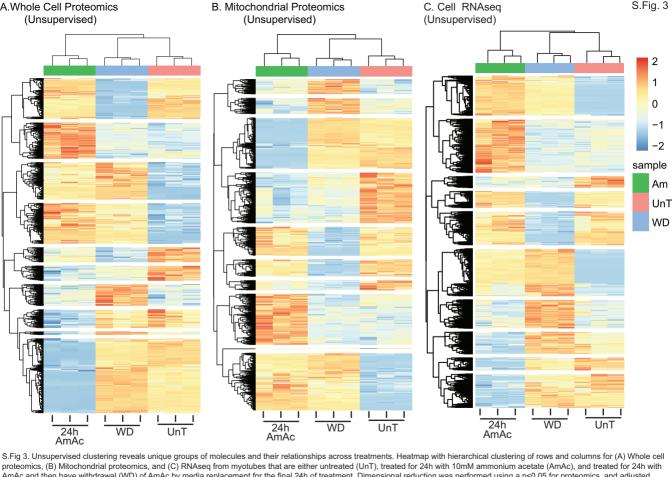
FCCP for measuring maximum respiration; 375 nM rotenone (Rot.) to inhibit Complex I; 125 nM antimycin A to inhibit Complex III; 2 mM ascorbate and 2 mM tetramethyl p-phenylene diamine to test complex IV activity; and 50 mM sodium azide to inhibit complex IV activity. R.R.; reserve respiratory capacity; Max. R.: maximum respiration. All data expressed as mean±SD from at least 4 sets of isolated mitochondria each group *p <0.05; **p <0.01; ***p <0.001 compared to untreated controls determined using unpaired Student's t-test.

S.Fig. 5. Ammonia rechallenge after withdrawal in C2C12 myotubes. (A) Representative photomicrographs and diameter of myotubes, expressed as a percentage of controls, that were either untreated, treated for 24h with ammonium acetate (AmAc), or following AmAc for 24h, ammonia withdrawal for 24h and subsequent rechallenge with AmAc for 24h (WD+Re.). Quantitative data are represented as box and whisker plots, with box bounds from 1st quartile to the 3rd quartile, median line, x as the mean, and whiskers ranging from minimum to maximum values, outliers represented as circles for myotube diameter as a percentage of controls. (B) Oxygen consumption measured by high resolution respirofluorometry in intact myotubes that have been treated with AmAc WD+Re. (C) ATP content of myotubes that were either untreated or treated with AmAc with or without WD+Re. (D) Representative immunoblots and densitometry for carbonylated proteins in untreated myotubes compared to those treated with AmAc or WD+Re. (E)

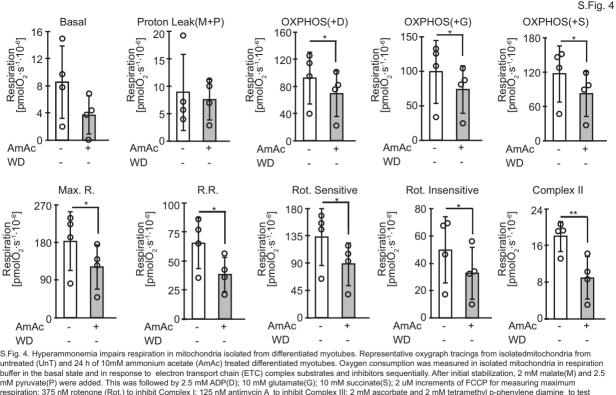
Representative immunoblots and densitometry for p16lNK and p21 in myotubes that were untreated, treated with AmAc or WD+Re. All data expressed as mean±SD from at least 6 biological replicates in oxygraph studies and at least 3 biological replicates in all other studies. *p<0.05; **p<0.01; ***p<0.001 using unpaired Student's t-test or one-way analysis of variance followed by Bonferroni post-hoc comparison tests.



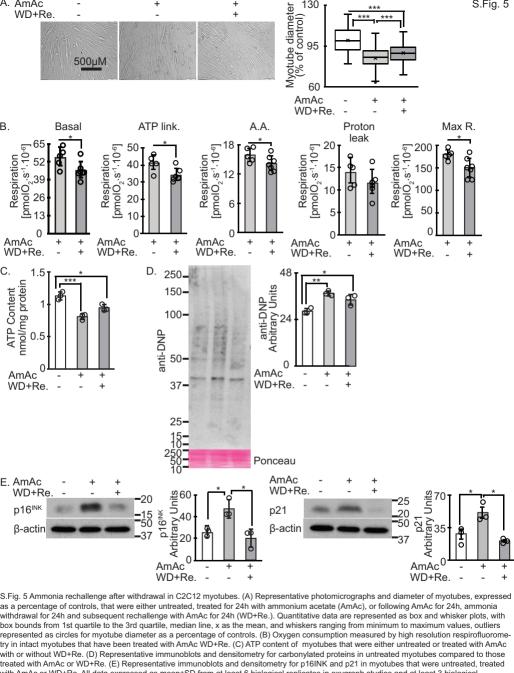




AmAc and then have withdrawal (WD) of AmAc by media replacement for the final 24h of treatment. Dimensional reduction was performed using a p<0.05 for proteomics, and adjusted p-value of 0.05 for RNAseq using unpaired Student's t-test following Benjamini Hochberg correction. All experiments were performed in n=3 biological replicates.



complex IV activity; and 50 mM sodium azide to inhibit complex IV activity. R.R.; reserve respiratory capacity; Max. R.: maximum respiration. All data expressed as mean±SD from at least 4 sets of isolated mitochondria each group *p<0.05; **p<0.01; ***p<0.001 compared to untreated controls determined using an unpaired Student's t-test.



130

S.Fig. 5

A. AmAc

with AmAc or WD+Re. All data expressed as mean±SD from at least 6 biological replicates in oxygraph studies and at least 3 biological replicates in all other studies. *p<0.05; **p<0.01; ***p<0.001 using unpaired Student's t-test or one-way analysis of variance followed by Bonferroni post-hoc comparison tests.

Supplementary Table 3. Electron transport chain proteins expressed in the cellular mitochondrial proteome.

Сарричина	AmAc vs WD vs WD vs				
Genes	UnT	AmAc	UnT	Complex	Complex I Module
MT-ND1	-2.37	N/A	-1.73	1	-
MT-ND4	-1.74	1.03	N/A	1	-
MT-ND5	-1.13	N/A	-0.68	1	-
NDUFA10	-1.32	N/A	-0.69	1	PP
NDUFA11	-1.16	N/A	-0.80	1	PP
NDUFA12	-1.32	N/A	-0.80	1	N/Q
NDUFA13	-1.06	N/A	-0.97	1	PP
NDUFA2	-0.92	N/A	-0.57	1	N
NDUFA4	-0.19	-0.20	-0.39	1	-
NDUFA5	-1.19	N/A	-0.63	1	Q
NDUFA6	-0.85	N/A	-0.59	1	Q
NDUFA7	-1.78	N/A	-1.37	1	Q
NDUFA8	-1.09	N/A	-0.73	1	PP
NDUFA9	-1.17	N/A	-0.83	1	Q
NDUFAB1	-0.45	N/A	-0.28	1	-
NDUFB10	-1.34	0.62	-0.71	1	PD
NDUFB11	-0.96	N/A	-0.67	1	PD
NDUFB4	-1.20	N/A	-0.84	1	PD
NDUFB5	-1.07	0.38	-0.69	1	PD
NDUFB6	-1.29	N/A	-0.97	1	PD
NDUFB7	-1.18	N/A	-0.72	1	PD
NDUFB8	-0.58	N/A	-0.48	1	PD
NDUFB9	-1.17	N/A	-0.78	1	PD
NDUFS1	-0.94	N/A	-0.50	1	N
NDUFS2	-1.09	N/A	-0.61	1	Q
NDUFS3	-1.04	N/A	-0.71	1	Q
NDUFS4	-1.19	N/A	-0.93	1	N
NDUFS6	-1.00	N/A	-0.82	1	N/Q
NDUFS7	-1.16	N/A	-0.66	1	Q
NDUFS8	-1.16	N/A	-0.73	1	Q
NDUFV1	-0.93	N/A	-0.60	1	N
NDUFV2	-0.69	N/A	-0.50	1	N
NDUFV3	-1.01	N/A	-0.79	1	PD
SDHA	-0.37	N/A	-0.36	2	-
SDHB	-0.50	N/A	-0.45	2	-
SDHD	-0.48	N/A	-0.52	2	-
CYC1	-0.73	N/A	-0.51	3	-
UQCR10	-0.71	N/A	-0.43	3	-
UQCRB	N/A	N/A	-0.64	3	-
UQCRC1	-0.97	N/A	-0.51	3	-
UQCRC2	-0.86	N/A	-0.46	3	-
UQCRQ	N/A	N/A	-0.52	3	-
COX15	-0.49	0.31	-0.19	4	-
COX4I1	-0.54	N/A	-0.42	4	-
COX5A	-0.57	0.15	-0.42	4	-
COX6A1	-1.53	N/A	-0.75	4	-
COX6B1	-1.17	N/A	-0.74	4	-
Cox6c	-1.21	N/A	-0.93	4	-
COX7A2	-1.00	N/A	-1.08	4	-
COX7A2L	-1.23	0.54	-0.70	4	-

CYB5A	N/A	N/A	-0.21	4	-
MT-CO2	-1.13	N/A	-0.91	4	-
SURF1	-1.44	0.95	-0.49	4	-
ATP5F1A	N/A	N/A	-0.15	5	-
ATP5F1B	-0.18	N/A	-0.14	5	-
ATP5F1C	N/A	N/A	-0.14	5	-
ATP5F1D	-0.12	N/A	-0.16	5	-
ATP5MF	-0.29	N/A	-0.24	5	-
ATP5MG	-0.23	N/A	-0.15	5	-
ATP5PD	-0.18	N/A	N/A	5	-
ATP5PF	N/A	N/A	-0.48	5	-
ATP5PO	N/A	N/A	-0.17	5	-
ATPAF2	N/A	-0.27	-0.29	5	-
CYCS	N/A	-0.32	-0.27	CytoChromeC	-

Numbers indicate log2 fold change of protein expression in isolated cellular mitochondrial untargeted quantitative proteomics; AmAc= ammonium acetate; UnT= untreated; WD= ammonia withdrawal; N/A = protein is not differentially expressed; N= dehydrogenase module (Complex I); N/Q= dehydrogenase module and/or hydrogenase module (Complex I); PD=; PP= proton transloacation module (Complex I); Q= hydrogenase module (Complex I); . p<0.05 in n=3 biological replicates.

REAGENT or RESOURCE	SOURCE	IDENTIFIER
Antibodies (clone number (experimental usage; dilu		
Mouse monoclonal anti-α-Tubulin (TU-02	Santa Cruz	Cat# sc-8035
(immunoblot normalization in isolated mitochondrial	Biotechnology,	
purity in UnT, AmAc, WD treated myotubes;	Dallas, Texas	
1:5000))		
Mouse monoclonal anti-β-Actin (C4 (immunoblot	Santa Cruz	Cat# sc-47778
normalization and senescence associated beta-	Biotechnology,	
galactosidase activity in UnT, AmAc, WD treated	Dallas, Texas	
myotubes and gastroc muscle Sham or PCA rats		
with or without LOLA-R; 1:5000))	D (' T	0.1//.40404
Mouse polyclonal anti-citrate synthase	ProteinTech,	Cat# 16131
(densitometry of CS in UnT, AmAc, WD treated	Rosemont, IL	
myotubes and gastroc muscle from Sham or PCA		
rats with or without LOLA-R; 1:5000)	Dathyd	Cot# 0.150 1170
Goat polyclonal anti-DNP (carbonylated in proteins	Bethyl	Cat# A150-117A
in UnT, AmAc, WD treated myotubes and in	Laboratories Inc.,	
gastroc muscle from Sham or PCA rats with or	Montgomery, TX	
without LOLA-R 1:10,000) Rabbit monoclonal anti-p16 INK4A (D7C1M)	Cell Signaling	Cat# 80772T
(senescence associated beta-galactosidase activity	Technology,	Cat# 007721
in UnT, AmAc, WD treated myotubes and gastroc	Danvers, MA	
muscle from Sham or PCA rats with or without;	Darivers, IVIA	
1:2000))		
Rabbit polyclonal anti-p21 (senescence associated	ProteinTech,	Cat# 10355-1-AP
beta-galactosidase activity in UnT, AmAc, WD	Rosemont, IL	Gath 10000 17 ti
treated myotubes and gastroc muscle from Sham	1 10001110111, 12	
or PCA gastroc; 1:2000))		
Mouse monoclonal anti-p53 (1C12 (senescence	Cell Signaling	Cat# 2524s
associated beta-galactosidase activity in UnT,	Technology,	
AmAc, WD treated myotubes; 1:2000))	Danvers, MA	
Rabbit polyclonal anti-pan-cadherin (densitometry	Cell Signaling	Cat# 4068
of isolated mitochondrial purity in UnT, AmAc, WD	Technology,	
treated myotubes; 1:2000)	Danvers, MA	
Mouse monoclonal anti-PDI (RL90 (densitometry of	Novus Biologicals,	Cat# NB300-517
isolated mitochondrial purity in UnT, AmAc, WD	Littleton, CO	
treated myotubes; 1:2000))		
Rabbit polyclonal anti-phospho-p53 (Ser15	Cell Signaling	Cat# 9284
(senescence associated beta-galactosidase activity	Technology,	
in UnT, AmAc, WD treated myotubes and gastroc	Danvers, MA	
muscle from Sham or PCA rat with or without		
LOLA-R; 1:1000))	0.411.01.555.5	C-# 4000
Rabbit polyclonal anti-VDAC (isolated	Cell Signaling	Cat# 4866
mitochondrial purity in UnT, AmAc, WD treated	Technology,	
myotubes; densitometry of voltage dependent	Danvers, MA	
anion channel in UnT, AmAc, WD treated		
myotubes and gastroc muscle from Sham or PCA rats with or without LOLA-R; 1:2000)		
Goat anti-mouse IgG3, fc gamma Specific, HRP	Cell Signaling	Cat# 75952
Conjugate (secondary antibody for anti-DNP;	Technology,	Oddir 10002
1:5000)	Danvers, MA	
Anti-rabbit IgG, HRP-Linked (secondary antibody	Cell Signaling	Cat# 7074s
for anti-p16 INK4, anti-p21, anti-phospho-p53, and	Technology,	000, 101 10
anti-pan-cadherin; 1:5000)	Danvers, MA	
and pair odditini, 1.0000/	Danvois, IVIA	

Anti-mouse IgG, HRP-Linked (secondary antibody	Cell Signaling	Cat# 7076s					
for anti-p53, anti-citrate synthase, anti-PDI, anti-α-	Technology,	Gain 10100					
Tubulin, anti-β-actin; 1:5000)	Danvers, MA						
Chemicals, peptides, and recombinant proteins							
α-Ketoglutaric acid	Sigma-Aldrich, St. Louis, MO	Cat# 75890					
α-Ketoglutaric acid- ¹³ C ₅	Cambridge Isotopes, Tewksbury, MA	Cat# CLM-2411-PK					
β-Mercaptoethanol	Sigma-Aldrich, St. Louis, MO	Cat# M3148					
(+)-Sodium L-ascorbate (Ascorbate)	Sigma-Aldrich, St. Louis, MO	Cat# A7631					
(L)-Malic Acid (Malate)	Sigma-Aldrich, St. Louis, MO	Cat# M1000					
2,4'-Dinitrophenylhydrazine (DNPH)	Sigma-Aldrich, St. Louis, MO	Cat# D199303					
2,7' -Dichloorofluorescin diacetate (DCFDA)	Sigma-Aldrich, St. Louis, MO	Cat# D6883					
3-[(3-cholamidopropyl) dimethylammonio]-1- propanesulfonate (CHAPS)	EMD Millipore Corp., Billerica, MA	Cat# 220201					
3,3' -Diaminobenzidine (DAB)	Sigma-Aldrich, St. Louis, MO	Cat# D8001					
4-Methlyumbeliferyl β-D-galactopyranoside (MUG)	Sigma-Aldrich, St. Louis, MO	Cat# M1633					
5-Bromo-4-Chloro-3-Indolyl β-D-Galactopyranoside (X-Gal)	ThermoFisher Scientific, Waltham, MA	Cat# B1690					
Adenosine diphosphate (ADP)	Sigma-Aldrich, St. Louis, MO	Cat# A5285					
Adenosine triphosphate (ATP)	Sigma-Aldrich, St. Louis, MO	Cat# A2383					
Ammonium acetate	Sigma-Aldrich, St. Louis, MO	Cat# A7330					
Antimycin a	Sigma-Aldrich, St. Louis, MO	Cat# A8674					
Benzamidine	Honeywell Fluka Chemicals, Charlotte, NC	Cat# 12072					
Bis-Tris	Sigma-Aldrich, St. Louis, MO	Cat# B4429					
Carbonyl cyanide p-trifluoro-methoxyphenyl hydrazone (FCCP)	Sigma-Aldrich, St. Louis, MO	Cat# C2920					
Citric acid	Sigma-Aldrich, St. Louis, MO	Cat# PHR1416					
Citric acid	Sigma-Aldrich, St. Louis, MO	Cat# 251275					
Citric acid- ¹³ C ₆	Cambridge Isotopes, Tewksbury, MA	Cat# CLM-9021-PK					
Coomassie brilliant stain G-250	Bio-Rad Laboratories, Hercules, CA	Cat# 1610406					

Coomassie brilliant stain R-250	Bio-Rad Laboratories, Hercules, CA	Cat# 1610400
Cytochrome c from bovine heart	Sigma-Aldrich, St. Louis, MO	Cat# C2037
Digitonin	Sigma-Aldrich, St. Louis, MO	Cat# D5628
Ethyl acetate	Fisher Scientific, Hampton. NH	Cat# E195SK-4
Fumaric acid	Sigma-Aldrich, St. Louis, MO	Cat# 47910
Fumaric acid- ¹³ C ₄	Cambridge Isotopes, Tewksbury, MA	Cat# CLM-1529-PK
Glutamate	Sigma-Aldrich, St. Louis, MO	Cat# G1626
Glycine	ThermoFisher Scientific, Waltham, MA	Cat# BP3815
L-Malic acid	Sigma-Aldrich, St. Louis, MO	Cat# 112577
L-Malic acid- ¹³ C ₄	Cambridge Isotopes, Tewksbury, MA	Cat# CLM-8065-PK
L-Ornithine L-Aspartate (LOLA)	Sigma-Aldrich, St. Louis, MO	Cat# 07125
Lead(II) nitrate	Sigma-Aldrich, St. Louis, MO	Cat# 228621
Magnesium chloride	Sigma-Aldrich, St. Louis, MO	Cat# 208337
Magnesium sulfate heptahydrate	Sigma-Aldrich, St. Louis, MO	Cat# M5921
MitoSOX™ red mitochondrial superoxide indicator	Invitrogen, ThermoFisher Scientific, Waltham, MA	Cat# M36008
N-Dodecyl β-D-maltoside 98%	Sigma-Aldrich, St. Louis, MO	Cat# D4641
N-tert-butyldimethylsilyl-N-methyltrifluroacetamide (MTBSTFA) and MTBSTFA + 1% TBDMCS (tert-butyldimethlysilyl ethers) sialylation reagent	ThermoFisher Scientific, Waltham, MA	Cat# TS-4890
N,N,N,N'-Tetramethyl-p-phenylenediamine dihydrochloride (TMPD)	Sigma-Aldrich, St. Louis, MO	Cat# T3134
NativePage™ 3 to 12%, bis-tris, 1.0 mm, mini gel	ThermoFisher Scientific, Waltham, MA	Cat# BN1001BOX
Nicotinamide adenine dinucleotide (NADH)	Sigma-Aldrich, St. Louis, MO	Cat# N6005
Nitro blue tetrazolium chloride (NBT)	ThermoFisher Scientific, Waltham, MA	Cat# N6495
Oligomycin	Sigma-Aldrich, St. Louis, MO	Cat# O4876
Phenylmethanesulfonylfluoride (PMSF)	Sigma-Aldrich, St. Louis, MO	Cat# P-7626

Potassium hexacyanoferrate (II) trihydrate	Sigma-Aldrich, St. Louis, MO	Cat# P3289			
Potassium hexacyanoferrate (III)	Sigma-Aldrich, St. Louis, MO	Cat# P8131			
Rifaximin	Sigma-Aldrich, St. Louis, MO	Cat# R9904			
Rotenone	Sigma-Aldrich, St. Louis, MO	Cat# R8875			
Sodium azide	Sigma-Aldrich, St. Louis, MO	Cat# S2002			
Sodium chloride	Fisher Scientific, Hampton. NH	Cat#S640			
Sodium phosphate	Sigma-Aldrich, St. Louis, MO	Cat# S5136			
Sodium pyruvate (Pyruvate)	Sigma-Aldrich, St. Louis, MO	Cat# P2256			
Sodium pyruvate- ¹³ C ₃	Cambridge	Cat# CLM-2440-PK			
Codium pyravato C ₃	Isotopes, Tewksbury, MA	Cath CLW 211011X			
Sodium succinate dibasic hexahydrate (Succinate)	Sigma-Aldrich, St. Louis, MO	Cat# S2378			
Succinic acid	Sigma-Aldrich, St.	Cat# 398055			
Succinic acid- ¹³ C ₄	Louis, MO Cambridge	Cat# CLM-1371-PK			
Succinic acid- C ₄	Isotopes,	Cat# CLIVI-1371-FK			
	Tewksbury, MA				
Tricine	Sigma-Aldrich, St. Louis, MO	Cat# T0377			
Critical Commercial Assays					
ATP determination kit	Invitrogen,	Cat# A22066			
	ThermoFisher				
	Scientific,				
	Waltham, MA				
Lipid peroxidation (MDA) assay kit	Abcam,	Cat# ab118970			
(Colorimetric/Fluorometric) (TBARS) MiR05-kit	Cambridge, UK O2k-Network Lab,	Cat# MiPNet22.10 MiR05-Kit			
MIROS-KIL	Innsbruck, Austria	Cat# MIPNet22.10 MIROS-Kit			
Deposited data					
Cellular proteomics	This paper	ProteomeXchange Consortium via			
Mitochondrial proteomics	This paper	the PRIDE partner repository with the dataset identifier PXD027754			
Tissue proteomics	This paper	and 10.6019/PXD027754			
Cellular RNAseq	This paper	https://github.com/dasaraslab/Unbia sed			
Experimental models: cell lines					
C2C12 myotubes	ATCC, Manassas, VA	Cat# CRL-1772			
Experimental models: organisms/strains					
Rats: Sprague-Dawley	Charles River	Strain code# 400			
	Laboratory,				
	Wilmington, MA				

Software and algorithms				
Adobe Illustrator 2021	Adobe, San Jose, CA	https://www.adobe.com/products/illu strator.html?sdid=KKQML&mv=sear ch&ef_id=EAlalQobChMlxbejkY7X8 wlVy3xvBB1b7whaEAAYASAAEgK RYfD_BwE:G:s&s_kwcid=AL!3085! 3!442365417815!e!!g!!adobe%20illu strator!1711729586!70905759510& gclid=EAlalQobChMlxbejkY7X8wlV y3xvBB1b7whaEAAYASAAEgKRYf D_BwE		
DatLab 6	Oroboros, Innsbruck, Austria	Cat# 27142-01		
g:Profiler	Open Source	https://biit.cs.ut.ee/gprofiler/gost		
ImageJ	NIH, Bethesda, MD	https://imagej.nih.gov/ij/		
IPA	Qiagen, Hilden, Germany	https://digitalinsights.qiagen.com/pro ducts-overview/discovery-insights- portfolio/analysis-and- visualization/qiagen-ipa/		
R studio	Open Source	https://www.rstudio.com/products/rst udio/download/		

Supplementary Table 7. Clusters of changes in expression of genes and proteins during hyperammonemia and following ammonia withdrawal/lowering

Cluster	Name	Expression change from untreated with AmAc treatment	Expression change from untreated with WD
Drogragaiya	a.	Increases	Increases further than the change with AmAc with WD
Progressive	j.	Decreases	Decreases further than the change with AmAc with WD
Persistent	b.	Increases	Increases with respect to UnT, but not significantly different compared to treatment with AmAc
reisistent	i.	Decreases	Decreases with respect to UnT, but not significantly different compared to treatment with AmAc
Dartially rayaread	d.	Increases	Increased with respect to UnT, but less than that seen with AmAc
Partially reversed	g.	Decreases	Decreased with respect to UnT, but less than that seen with AmAc
Completely	e.	Increases	No significant difference compared to
Reversed	f.	Decreases	UnT
Overcorrection	C.	Increases	Decreases
Overcorrection	h.	Decreases	Increases

AmAc: 24h 10mM ammonium acetate treatment; UnT: untreated; WD: Treatment with 24h 10mM ammonium acetate followed by 24h ammonium acetate withdrawal

Supplementary Table 8. Verified and non-verified mitochondrial proteins against MitoCarta3.0

			Verified		
	Whole Cell	Mito	MitoCarta		
	proteome	proteome	3.0	Count	Rationale
					All verified mitochondrial proteins found in the
Group 1	yes	yes	yes	356	mitochondria also detected in the whole cell
					Mitochondrial targeted proteins detected in the
Group 2	yes	no	yes	22	whole cell that did not enter the mitochondria
					Mitochondrial proteins that are not
					concentrated enough to be present in the
Group 3	no	yes	yes	170	whole cell lysate
					Verified mitochondrial proteins not detected in
Group 4	no	no	yes	592	C2C12 cells or mitochondria
Group 5	yes	no	no	634	Non mitochondrial whole cell proteins
					Non verified mitochondrial proteins OR
Group 6	no	yes	no	400	transported proteins
					Transported cytosolic or other non-
					mitochondrial proteins OR non-verified
Group 7	yes	yes	no	1197	mitochondrial proteins

Mito: isolated mitochondrial proteome